## What is claim d:

- 1. A receptor comprised of at least one T1R1 polypeptide (or a variant, fragment, or chimera of said T1R1 polypeptide) and/or at least one T1R3 polypeptide (or a variant, fragment, or chimera of said T1R3 polypeptide), wherein said receptor specifically binds to and/or is activated by umami taste stimuli.
- 2. The receptor of Claim 1 containing a fragment, variant, or chimera of a native hT1R1 polypeptide.
- 3. The receptor of Claim 1 containing a fragment, variant, or chimera of a native hT1R3 polypeptide.
- 4. The receptor of Claim 1 wherein said T1R1 and T1R3 are derived from the same species.
- 5. The receptor of Claim 1 wherein said T1R1 and T1R3 are derived from different species.
- 6. The receptor of Claim 1 wherein said T1R1 and/or T1R3 are derived from a mammal, fish, reptile, amphibian, or bird.
- 7. The receptor of Claim 1 wherein said T1R1 and/or T1R3 are from the group consisting of: hT1R1, hT1R3, rT1R1, rT1R3, mT1R1, mT1R3; or fragments, variants, or chimeras derived therefrom.
- 8. A composition that contains a receptor according to Claim 1.
- 9. A composition that contains a receptor according to Claim 2.

- 10. A composition that contains a receptor according to Claim 3.
- 11. A composition that contains a receptor according to Claim 4.
- 12. A composition that contains a receptor according to Claim 5.
- 13. A composition that contains a receptor according to Claim 6.
- 14. A composition that contains a receptor according to Claim 7.
- 15. A cell that expresses a receptor comprised of at least one T1R1 polypeptide (or a variant, fragment, or chimera of said T1R1 polypeptide) and/or at least one T1R3 polypeptide (or a variant, fragment, or chimera of said T1R3 polypeptide), wherein said receptor specifically binds to and/or is activated by umami taste stimuli.
- 16. The cell of Claim 15, which is selected from the group consisting of HEK-293, COS and CHO cells, and Xenopus oocytes.
- 17. The cell of Claim 15 wherein said T1R1 and T1R3 are of the same species.
- 18. The cell of Claim 15 wherein said T1R1 and T1R3 are derived from different species.
- 19. The cell of Claim 15 wherein said T1R1 and/or T1R3 are derived from a mammal, fish, reptile, amphibian, or bird.

- The cell of Claim 15 wherein said T1R1 and T1R3 are selected from the group consisting of: hT1R1, hT1R3, mT1R1, mT1R3, rT1R1 and rT1R3; or fragments, variants or chimeras therefrom.
- A receptor comprised of at least one T1R2 polypeptide (or a variant, fragment, or chimera of said T1R2 polypeptide) and/or at least one T1R3 polypeptide (or a variant, fragment, or chimera of said T1R3 polypeptide), wherein said receptor specifically binds to and/or is activated by sweet taste stimuli.
- 22. The receptor of Claim 21 containing a fragment, variant, or chimera of a native hT1R2 polypeptide.
- 23. The receptor of Claim 21 containing a fragment, variant, or chimera of a native hT1R3 polypeptide.
- 24. The receptor of Claim 21 wherein said T1R2 and T1R3 are derived from the same species.
- 25. The receptor of Claim 21 wherein said T1R2 and T1R3 are derived from different species.
- 26. The receptor of Claim 21 wherein said T1R2 and/or T1R3 are derived from a mammal, fish, reptile, amphibian, or bird.
- The receptor of Claim 21 wherein said T1R2 and/or T1R3 are from the group consisting of: hT1R2, hT1R3, rT1R2, rT1R3, mT1R2, mT1R3; or fragments, variants, or chimeras derived therefrom.
- 28. A composition that contains a receptor according to Claim 21.
- A composition that contains a receptor according to Claim 22.

- 30. A composition that contains a receptor according to Claim 23.
- 31. A composition that contains a receptor according to Claim 24.
- 32. A composition that contains a receptor according to Claim 25.
- 33. A composition that contains a receptor according to Claim 26.
- 34. A composition that contains a receptor according to Claim 27.
- 35. A cell that expresses a receptor comprised of at least one T1R2 polypeptide (or a variant, fragment, or chimera of said T1R2 polypeptide) and/or at least one T1R3 polypeptide (or a variant, fragment, or chimera of said T1R3 polypeptide), wherein said receptor specifically binds to and/or is activated by sweet taste stimuli.
- 36. The cell of Claim 35, which is selected from the group consisting of HEK-293, COS and CHO cells, and Xenopus oocytes.
- 37. The cell of Claim 35 wherein said T1R2 and T1R3 are of the same species.
- 38. The cell of Claim 35 wherein said T1R2 and T1R3 are derived from different species.
- 39. The cell of Claim 35 wherein said T1R2 and/or T1R3 are derived from a mammal, fish, reptile, amphibian, or bird.

- The cell of Claim 35 wherein said T1R2 and T1R3 are selected from the group consisting of: hT1R2, hT1R3, mT1R2, mT1R3, rT1R2 and rT1R3; or fragments, variants or chimeras therefrom.
- 41. A receptor of Claims 1-7 or 21-27 bound to a solid phase.
- 42. A receptor of Claims 1-7 or 21-27 in solution.
- 43. A receptor of Claims 1-7 or 21-27 in a lipid bilayer or vesicle.
- A method for identifying compounds that modulate taste perception by identifying compounds that bind to, activate, inhibit, enhance and/or modulate one or more of the receptors of Claims 1-7 and/or 21-27.
- 45. The method of Claim 44 wherein the receptor is bound to a solid phase.
- 46. The method of Claim 44 wherein the receptor is in solution.
- The method of Claim 44 wherein the receptor is in a lipid bilayer or vesicle.
- 48. The method of Claim 44 which uses a receptor-binding assay to identify the said compound.
- The method of Claim 44 which uses a receptor activity-based assay to identify the said compound.
- 50. The method of Claim 44 wherein said receptor is expressed in a cell.

- 51. The method of Claim 50 wherein said cell is a HEK-293, COS, or CHO cell.
- 52. The method of Claim 50 wherein said compound is identified by its effect on receptor internalization.
- 53. The method of Claim 50 wherein said compound is identified by its effect on receptor phosphorylation.
- 54. The method of Claim 50 wherein said compound is identified by its effect on arrestin translocation.
- 55. The method of Claim 44 which uses an assay for second messengers.
- 56. The method of Claim 55 wherein the said second messenger is cAMP or IP<sub>3</sub>.
- 57. The method of Claim 50 which uses a voltage-sensitive or calciumsensitive dye.
- 58. The method of Claim 50 wherein said cell expresses at least one G protein.
- 59. The method of Claim 58 wherein said G protein is a promiscuous G protein.
- 60. The method of Claim 59 wherein said promiscuous G protein is  $G_{\sigma 15}$  or  $G_{\sigma 16}$ .

- The method of Claim 44 wherein said receptor is expressed using a viral vector.
- 62. The method of Claim 61 wherein said viral vector is expressed in a mammalian cell.
- 63. The method of Claim 44 wherein said receptor is produced by in vitro translation.
- 64. The method of Claim 44 wherein said receptor is isolated in a membrane-bound form.
- The method of Claim 44 wherein said compound is identified by its effect on G protein activation by said receptor.
- The method of Claim 44 wherein said compound is identified by its effect on receptor conformation.
- 67. The method of Claim 66 wherein said conformation change is detected by altered susceptibility to proteolysis.
- 68. The method of Claim 66 wherein said conformation change is detected by NMR spectroscopy.
- 69. The method of Claim 66 wherein said conformation change is detected by fluorescence spectroscopy.
- 70. The method of Claim 44 wherein said compound is identified by its effect on binding of a radioactively or fluorescently labeled ligand to said receptor.

- 71. The method of Claim 70 wherein displacement of said labeled compound is determined by fluorescence polarization or FRET assay.
- 72. The method of Claim 44 which is a high-throughput screening assay.
- 73. The method of Claim 44 wherein receptor activity is linked to a reporter gene.
- 74. The method of Claim 73 wherein said reporter gene is luciferase, alkaline phosphatase,  $\beta$ -galactosidase, or  $\beta$ -lactamase.
- 75. The method of Claim 44 wherein said receptor is a constitutively active variant.
- 76. The method of Claim 44 wherein expression of said receptor is under the control of a constitutive promoter.
- 77. The method of Claim 44 wherein expression of said receptor is under the control of a regulated promoter.
- 78. The method of Claim 44 wherein said receptor is fused to a peptide that facilitates surface expression.
- 79. The method of Claim 78 wherein said peptide is a PDZ-domain-interacting peptide.
- 80. The method of Claim 44 wherein the effect of said compound on said receptor is predicted based on the X-ray crystal structure of said receptor.

- The method of Claim 44 wherein said compound is identified by its effect on a non-human animal expressing native or transgenic T1R receptors.
- 82. The method of Claim 81 wherein said compound is identified by its effect on behavior.
- 83. The method of Claim 81 wherein said compound is identified by its effect on taste receptor cells.
- 84. The method of Claim 81 wherein said non-human animal is a mouse, rat, worm, fish, or insect.
- 85. The method of Claim 44 wherein said compound is identified by its effect on a yeast cell expressing said receptor.
- 86. The method of Claim 44 wherein said compound is identified from a combinatorial library of compounds.
- 87. The method of Claim 44 wherein said compound is identified from a peptide library.
- 88. The method of Claim 44 wherein said compound is identified from a randomized library of small molecules.
- 89. A method of modifying taste sensation in an animal using compounds identified according to Claim 44.
- 90. The method of Claim 89 wherein said taste sensation is umami taste.
- 91. The method of Claim 89 wherein said taste sensation is sweet taste.

- 92. The method of Claim 89 wherein said animal is a human, dog, cat, fish, cow, sheep, or pig.
- 93. The method of Claim 89 wherein said compound is formulated in a food, beverage, or oral pharmaceutical composition.
- 94. A method of quantifying the taste of individual compounds or food or beverage compositions using one or more of the receptors of Claims 1-7 and/or 21-27.
- 95. A cell that stably expresses a receptor comprised of at least one T1R1 polypeptide or a variant, fragment, or chimera of said T1R1 polypeptide and/or at least one T1R3 polypeptide or a variant, fragment, or chimera of said T1R3 polypeptide, wherein said receptor specifically binds to and/or is activated by umami taste stimuli.
- 96. The cell of Claim 95, which is selected from the group consisting of HEK-293, COS and CHO cells, and Xenopus oocytes.
- 97. The cell of Claim 95 wherein said T1R1 and T1R3 are of the same species.
- 98. The cell of Claim 95 wherein said T1R1 and T1R3 are derived from different species.
- 99. The cell of Claim 95 wherein said T1R1 and/or T1R3 are derived from a mammal, fish, reptile, amphibian, or bird.

- The cell of Claim 95 wherein said T1R1 and T1R3 are selected from the group consisting of: hT1R1, hT1R3, mT1R1, mT1R3, rT1R1 and rT1R3; or fragments, variants or chimeras thereof.
- 101. The cell of Claim 95 which is an HEK-293 cell that stably expresses  $G_{\alpha 15}$ .
- 102. A composition that contains a cell according to Claim 95.
- 103. A composition that contains a cell according to Claim 96.
- 104. A composition that contains a cell according to Claim 97.
- 105. A composition that contains a cell according to Claim 98.
- 106. A composition that contains a cell according to Claim 99.
- 107. A composition that contains a cell according to Claim 100.
- 108. A composition that contains a cell according to Claim 101.
- 109. A cell that stably expresses a receptor comprised of at least one T1R2 polypeptide or a variant, fragment, or chimera of said T1R2 polypeptide and/or at least one T1R3 polypeptide or a variant, fragment, or chimera of said T1R3 polypeptide, wherein said receptor specifically binds to and/or is activated by sweet taste stimuli.
- 110. The cell of Claim 109, which is selected from the group consisting of HEK-293, COS and CHO cells, and Xenopus oocytes.

- 111. The cell of Claim 109 wherein said T1R2 and T1R3 are of the same species.
- 112. The cell of Claim 109 wherein said T1R2 and T1R3 are derived from different species.
- 113. The cell of Claim 109 wherein said T1R2 and/or T1R3 are derived from a mammal, fish, reptile, amphibian, or bird.
- The cell of Claim 109 wherein said T1R2 and T1R3 are selected from the group consisting of: hT1R2, hT1R3, mT1R2, mT1R3, rT1R2 and rT1R3; or fragments, variants or chimeras therefrom.
- 115. The cell of Claim 109 which stably expresses hT1R2, hT1R3 and a  $G_{\alpha 15}$ .
- 116. A method for identifying compounds that modulate taste perception by identifying compounds that bind to, activate, inhibit, and/or modulate a receptor expressed by a cell that stably expresses at least one T1R.
- 117. The method of Claim 116 wherein the cell is bound to a solid phase.
- 118. The method of Claim 117 wherein the cell is in solution.
- 119. The method of Claim 117 which uses a receptor-binding assay to identify the said compound.
- The method of Claims 116 wherein said cell is according to any one of Claims 95-101 or 109-115.

- 121. The method of Claim 116 which uses a receptor activity-based assay to identify the said compound.
- 122. The method of Claim 116 wherein said T1R receptor is expressed in a cell.
- 123. The method of Claim 116 wherein said cell is a HEK-293, COS, or CHO cell.
- 124. The method of Claim 116 wherein said compound is identified by its effect on receptor internalization by said cell.
- 125. The method of Claim 116 wherein said compound is identified by its effect on receptor phosphorylation.
- 126. The method of Claim 116 wherein said compound is identified by its effect on arresting translocation.
- 127. The method of Claim 116 which uses an assay for second messengers.
- 128. The method of Claim 127 wherein the said second messenger is cAMP or IP<sub>3</sub>.
- 129. The method of Claim 116 which uses a voltage-sensitive or calciumsensitive dye.
- 130. The method of Claim 116 wherein said cell expresses at least one G protein.

- 131. The method of Claim 130 wherein said G protein is a promiscuous G protein.
- The method of Claim 131 wherein said promiscuous G protein is  $G_{\sigma 15}$  or  $G_{\sigma 16}$ .
- 133. The method of Claim 116 wherein said receptor is stably expressed using a viral promoter.
- 134. The method of Claim 116 wherein said cell is a mammalian cell.
- 135. The method of Claim 116 wherein said compound is identified by its effect on G protein activation by said receptor.
- 136. The method of Claim 116 wherein said compound is identified by its effect on receptor conformation.
- 137. The method of Claim 136 wherein said conformation change is detected by altered susceptibility to proteolysis.
- 138. The method of Claim 136 wherein said conformation change is detected by NMR spectroscopy.
- 139. The method of Claim 136 wherein said conformation change is detected by fluorescence spectroscopy.
- 140. The method of Claim 136 wherein said compound is identified by its effect on binding of a radioactively or fluorescently labeled ligand to said stably expressed receptor.

- 141. The method of Claim 140 wherein displacement of said labeled compound is determined by fluorescence polarization or FRET assay.
- 142. The method of Claim 116 which is a high-throughput screening assay.
- 143. The method of Claim 116 wherein receptor activity is linked to a reporter gene.
- 144. The method of Claim 143 wherein said reporter gene is luciferase, alkaline phosphatase,  $\beta$ -galactosidase, or  $\beta$ -lactamase.
- 145. The method of Claim 140 wherein said receptor is a constitutively active variant.
- 146. The method of Claim 140 wherein expression of said receptor is under the control of a constitutive promoter.
- 147. The method of Claim 140 wherein expression of said receptor is under the control of a regulated promoter.
- 148. The method of Claim 140 wherein said receptor is fused to a peptide that facilitates surface expression.
- 149. The method of Claim 140 wherein said peptide is a PDZ-domain-interacting peptide.
- 150. The method of Claim 140 wherein the effect of said compound on said receptor is predicted based on the X-ray crystal structure of said receptor.

- 151. The method of Claim 116 wherein said compound is identified by its effect on a yeast cell expressing said receptor.
- 152. The method of Claim 116 wherein said compound is identified from a combinatorial library of compounds.
- 153. The method of Claim 116 wherein said compound is identified from a peptide library.
- 154. The method of Claim 116 wherein said compound is identified from a randomized library of small molecules:
- 155. A method of modifying taste sensation in an animal using compounds identified according to Claim 116.
- 156. The method of Claim 155 wherein said taste sensation is umami taste.
- 157. The method of Claim 155 wherein said taste sensation is sweet taste.
- 158. The method of Claim 155 wherein said animal is a human, dog, cat, fish, cow, sheep, or pig.
- 159. The method of Claim 155 wherein said compound is formulated in a food, beverage, or oral pharmaceutical composition.
- 160. A method of quantifying the taste of individual compounds or food or beverage compositions using a cell that stably expresses a heterologous nucleic acid sequence encoding at least one T1R according to one of Claims 1-7 or 15-21.

- 161. A cell line which inducibly expresses the human T1R1/T1R3 umami taste receptor or the T1R2/T1R3 sweet taste receptor.
- 162. The cell line of claim 161 which the cell line is a CHO, COS, HEK or BHK cell line.
- 163. The cell line of claim 162 which is an HEK-293 cell line.
- 164. The cell line of claim 161 which expresses a G protein.
- 165. The cell line of claim 164 wherein said G protein in Ga<sub>15</sub> or Ga<sub>16</sub>.
- 166. The cell line of claim 161 which stably expresses said T1R1/T1R3 receptor.
- 167. The cell line of claim 161 wherein the expression is induced by the GeneSwitch protein.
- 168. A method of using the cell line of claim 161 to identify a compound that agonizes or antagonizes the T1R1/T1R3 receptor or T1R2/T1R3 receptor.
- 169. The method of claim 168 which is a binding assay.
- 170. The method of claim 168 which is a high throughput screening assay.
- 171. The method of claim 168 which is fluorometric assay.
- 172. The method of claim 168 which screens for a compound that competes with L-glutamate or L-aspartate for binding to the T1R1/T1R3 umami taste receptor.

- 173. The method of claim 168 which is a high throughput screening assay that uses automated fluorometric imaging instrumentation.
- 174. The method of claim 168 which is used to screen a compound library for compounds that enhance or modulate the activity of L-glutamate to activate the T1R1/T1R3 umami taste receptor.
- 175. The method of claim 168 which is used to screen a compound library for compounds that agonize or antagonize the T1R2/T1R3 sweet taste receptor.
- 176. The method of claim 168 which screens for a compound that competes with IMP, GMP or their analogues for binding to the T1R1/T1R3 umami taste receptor.
- 177. The method of claim 168 which is used to screen a compound library for compounds that mimic the activity of IMP, GMP or their analogues that enhance the activity of a T1R1/T1R3 agonist.
- 178. The method of claim 168 which is used to screen a compound library for compounds that enhance or modulate the activity of a sweetener to activate the T1R2/T1R3 sweet taste receptor.
- 179. A method of inhibiting the T1R1/T1R3 umami taste receptor comprising contacting said receptor with a sweet-taste inhibitor that also inhibits both the T1R1/T1R3 sweet taste receptor and the T1R2/T1R3 taste receptor.
  - 180. The method of claim 179 wherein said inhibitor is lactisole.
- 181. A method for identifying compounds that modulate the T1R1/T1R3 umami taste receptor by screening for compounds that compete with lactisole for binding to and/or inhibiting the T1R1/T1R3 umami taste receptor.

- . 182. The method of claim 179 which is a cell-based assay.
- 183. The method of claim 182 wherein said assay uses a cell line that coexpresses T1R1 and T1R3.
- 184. The method of claim 183 wherein said cell line is a HEK-G $\alpha_{15}$  cell line.
- 185. The method of claim 183 wherein said cell line stably expresses said receptors.
- 186. The method of claim 183 wherein said cell line transiently expresses said receptors.
  - 187. The method of claim 186 wherein said G-protein is  $G\alpha_{15}$  or  $G\alpha_{16}$ .
  - 188. The method of claim 181 which is a cell-based assay.
- 189. The method of claim 188 wherein said assay uses a cell line that coexpresses T1R1 and T1R3.
- 190. The method of claim 189 wherein said cell line is a HEK-G $\alpha_{15}$  said cell line.
- 191. The method of claim 189 wherein said cell line stably expresses said receptors.
- 192. The method of claim 191 wherein said cell line transiently expresses said receptors.

193. The method of claim 192 wherein said G-protein is  $G\alpha_{15}$  or  $G\alpha_{16}$ .